

OIKE

RAW SEQUENCE LISTING

DATE: 04/23/2001

PATENT APPLICATION: US/09/821,883

TIME: 13:12:41

Input Set : A:\7636-0022.30-SEQLIST.txt

Output Set: N:\CRF3\04232001\I821883.raw

4 <110> APPLICANT: Laus, Reiner
5 Vidovic, Damir
6 Graddis, Thomas
8 <120> TITLE OF INVENTION: Compositions and Methods for Dendritic
9 Cell-Based Immunotherapy
12 <130> FILE REFERENCE: 7636-0022.30
C--> 14 <140> CURRENT APPLICATION NUMBER: US/09/821,883
C--> 15 <141> CURRENT FILING DATE: 2001-03-30
17 <150> PRIOR APPLICATION NUMBER: US 60/193,504
18 <151> PRIOR FILING DATE: 2000-03-30
20 <160> NUMBER OF SEQ ID NOS: 30
22 <170> SOFTWARE: FastSEQ for Windows Version 4.0
24 <210> SEQ ID NO: 1
25 <211> LENGTH: 555
26 <212> TYPE: PRT
27 <213> ORGANISM: Artificial Sequence
29 <220> FEATURE:
30 <223> OTHER INFORMATION: HER500 construct
32 <400> SEQUENCE: 1
33 Met Arg Ala Ala Pro Leu Leu Leu Ala Arg Ala Ala Ser Leu Ser Leu
34 1 5 10 15
35 Gly Phe Leu Phe Leu Leu Phe Phe Trp Leu Asp Arg Ser Val Leu Ala
36 20 25 30
37 Lys Glu Leu Ala Arg Gly Ala Ala Ser Thr Gln Val Cys Thr Gly Thr
38 35 40 45
39 Asp Met Lys Leu Arg Leu Pro Ala Ser Pro Glu Thr His Leu Asp Met
40 50 55 60
41 Leu Arg His Leu Tyr Gln Gly Cys Gln Val Val Gln Gly Asn Leu Glu
42 65 70 75 80
43 Leu Thr Tyr Leu Pro Thr Asn Ala Ser Leu Ser Phe Leu Gln Asp Ile
44 85 90 95
45 Gln Glu Val Gln Gly Tyr Val Leu Ile Ala His Asn Gln Val Arg Gln
46 100 105 110
47 Val Pro Leu Gln Arg Leu Arg Ile Val Arg Gly Thr Gln Leu Phe Glu
48 115 120 125
49 Asp Asn Tyr Ala Leu Ala Val Leu Asp Asn Gly Asp Pro Leu Asn Asn
50 130 135 140
51 Thr Thr Pro Val Thr Gly Ala Ser Pro Gly Gly Leu Arg Glu Leu Gln
52 145 150 155 160
53 Leu Arg Ser Leu Thr Glu Ile Leu Lys Gly Gly Val Leu Ile Gln Arg
54 165 170 175
55 Asn Pro Gln Leu Cys Tyr Gln Asp Thr Ile Leu Trp Lys Asp Ile Phe
56 180 185 190
57 His Lys Asn Asn Gln Leu Ala Leu Thr Leu Ile Asp Thr Asn Arg Ser
58 195 200 205
59 Arg Ala Cys His Pro Cys Ser Pro Met Cys Lys Gly Ser Arg Cys Trp
60 210 215 220

ENTERED

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61 Gly Glu Ser Ser Glu Asp Cys Gln Ser Leu Thr Arg Thr Val Cys Ala
62 225                230                235                240
63 Gly Gly Cys Ala Arg Cys Lys Gly Pro Leu Pro Thr Asp Cys Cys His
64                245                250                255
65 Glu Gln Cys Ala Ala Gly Cys Thr Gly Pro Lys His Ser Asp Cys Leu
66                260                265                270
67 Ala Cys Leu His Phe Asn His Ser Gly Ile Cys Glu Leu His Cys Pro
68                275                280                285
69 Ala Leu Val Thr Tyr Asn Thr Asp Thr Phe Glu Ser Met Pro Asn Pro
70                290                295                300
71 Glu Gly Arg Tyr Thr Phe Gly Ala Ser Cys Val Thr Ala Cys Pro Tyr
72 305                310                315                320
73 Asn Tyr Leu Ser Thr Asp Val Gly Ser Gly Ala Gly Gly Met Val His
74                325                330                335
75 His Arg His Arg Ser Ser Ser Thr Arg Ser Gly Gly Gly Asp Leu Thr
76                340                345                350
77 Leu Gly Leu Glu Pro Ser Glu Glu Ala Pro Arg Ser Pro Leu Ala
78                355                360                365
79 Pro Ser Glu Gly Ala Gly Ser Asp Val Phe Asp Gly Asp Leu Gly Met
80                370                375                380
81 Gly Ala Ala Lys Gly Leu Gln Ser Leu Pro Thr His Asp Pro Ser Pro
82 385                390                395                400
83 Leu Gln Arg Tyr Ser Glu Asp Pro Thr Val Pro Leu Pro Ser Glu Thr
84                405                410                415
85 Asp Gly Tyr Val Ala Pro Leu Thr Cys Ser Pro Gln Pro Glu Tyr Val
86                420                425                430
87 Asn Gln Pro Asp Val Arg Pro Gln Pro Pro Ser Pro Arg Glu Gly Pro
88                435                440                445
89 Leu Pro Ala Ala Arg Pro Ala Gly Ala Thr Leu Glu Arg Ala Lys Thr
90                450                455                460
91 Leu Ser Pro Gly Lys Asn Gly Val Val Lys Asp Val Phe Ala Phe Gly
92 465                470                475                480
93 Gly Ala Val Glu Asn Pro Glu Tyr Leu Thr Pro Gln Gly Gly Ala Ala
94                485                490                495
95 Pro Gln Pro His Pro Pro Pro Ala Phe Ser Pro Ala Phe Asp Asn Leu
96                500                505                510
97 Tyr Tyr Trp Asp Gln Asp Pro Pro Glu Arg Gly Ala Pro Pro Ser Thr
98                515                520                525
99 Phe Lys Gly Thr Pro Thr Ala Glu Asn Pro Glu Tyr Leu Gly Leu Asp
100                530                535                540
101 Val Pro Ala Ala Ala His His His His His
102 545                550                555
104 <210> SEQ ID NO: 2
105 <211> LENGTH: 690
106 <212> TYPE: PRT
107 <213> ORGANISM: Artificial Sequence
109 <220> FEATURE:
110 <223> OTHER INFORMATION: HER500-hGM-CSF construct
112 <400> SEQUENCE: 2

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113 Met Arg Ala Ala Pro Leu Leu Leu Ala Arg Ala Ala Ser Leu Ser Leu
114 1 5 10 15
115 Gly Phe Leu Phe Leu Leu Phe Phe Trp Leu Asp Arg Ser Val Leu Ala
116 20 25 30
117 Lys Glu Leu Ala Arg Gly Ala Ala Ser Thr Gln Val Cys Thr Gly Thr
118 35 40 45
119 Asp Met Lys Leu Arg Leu Pro Ala Ser Pro Glu Thr His Leu Asp Met
120 50 55 60
121 Leu Arg His Leu Tyr Gln Gly Cys Gln Val Val Gln Gly Asn Leu Glu
122 65 70 75 80
123 Leu Thr Tyr Leu Pro Thr Asn Ala Ser Leu Ser Phe Leu Gln Asp Ile
124 85 90 95
125 Gln Glu Val Gln Gly Tyr Val Leu Ile Ala His Asn Gln Val Arg Gln
126 100 105 110
127 Val Pro Leu Gln Arg Leu Arg Ile Val Arg Gly Thr Gln Leu Phe Glu
128 115 120 125
129 Asp Asn Tyr Ala Leu Ala Val Leu Asp Asn Gly Asp Pro Leu Asn Asn
130 130 135 140
131 Thr Thr Pro Val Thr Gly Ala Ser Pro Gly Gly Leu Arg Glu Leu Gln
132 145 150 155 160
133 Leu Arg Ser Leu Thr Glu Ile Leu Lys Gly Gly Val Leu Ile Gln Arg
134 165 170 175
135 Asn Pro Gln Leu Cys Tyr Gln Asp Thr Ile Leu Trp Lys Asp Ile Phe
136 180 185 190
137 His Lys Asn Asn Gln Leu Ala Leu Thr Leu Ile Asp Thr Asn Arg Ser
138 195 200 205
139 Arg Ala Cys His Pro Cys Ser Pro Met Cys Lys Gly Ser Arg Cys Trp
140 210 215 220
141 Gly Glu Ser Ser Glu Asp Cys Gln Ser Leu Thr Arg Thr Val Cys Ala
142 225 230 235 240
143 Gly Gly Cys Ala Arg Cys Lys Gly Pro Leu Pro Thr Asp Cys Cys His
144 245 250 255
145 Glu Gln Cys Ala Ala Gly Cys Thr Gly Pro Lys His Ser Asp Cys Leu
146 260 265 270
147 Ala Cys Leu His Phe Asn His Ser Gly Ile Cys Glu Leu His Cys Pro
148 275 280 285
149 Ala Leu Val Thr Tyr Asn Thr Asp Thr Phe Glu Ser Met Pro Asn Pro
150 290 295 300
151 Glu Gly Arg Tyr Thr Phe Gly Ala Ser Cys Val Thr Ala Cys Pro Tyr
152 305 310 315 320
153 Asn Tyr Leu Ser Thr Asp Val Gly Ser Gly Ala Gly Gly Met Val His
154 325 330 335
155 His Arg His Arg Ser Ser Ser Thr Arg Ser Gly Gly Gly Asp Leu Thr
156 340 345 350
157 Leu Gly Leu Glu Pro Ser Glu Glu Glu Ala Pro Arg Ser Pro Leu Ala
158 355 360 365
159 Pro Ser Glu Gly Ala Gly Ser Asp Val Phe Asp Gly Asp Leu Gly Met
160 370 375 380
161 Gly Ala Ala Lys Gly Leu Gln Ser Leu Pro Thr His Asp Pro Ser Pro

```

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```

162 385          390          395          400
163 Leu Gln Arg Tyr Ser Glu Asp Pro Thr Val Pro Leu Pro Ser Glu Thr
164          405          410          415
165 Asp Gly Tyr Val Ala Pro Leu Thr Cys Ser Pro Gln Pro Glu Tyr Val
166          420          425          430
167 Asn Gln Pro Asp Val Arg Pro Gln Pro Pro Ser Pro Arg Glu Gly Pro
168          435          440          445
169 Leu Pro Ala Ala Arg Pro Ala Gly Ala Thr Leu Glu Arg Ala Lys Thr
170          450          455          460
171 Leu Ser Pro Gly Lys Asn Gly Val Val Lys Asp Val Phe Ala Phe Gly
172 465          470          475          480
173 Gly Ala Val Glu Asn Pro Glu Tyr Leu Thr Pro Gln Gly Gly Ala Ala
174          485          490          495
175 Pro Gln Pro His Pro Pro Pro Ala Phe Ser Pro Ala Phe Asp Asn Leu
176          500          505          510
177 Tyr Tyr Trp Asp Gln Asp Pro Pro Glu Arg Gly Ala Pro Pro Ser Thr
178          515          520          525
179 Phe Lys Gly Thr Pro Thr Ala Glu Asn Pro Glu Tyr Leu Gly Leu Asp
180          530          535          540
181 Val Pro Ala Ala Ala Pro Ala Arg Ser Pro Ser Pro Ser Thr Gln Pro
182 545          550          555          560
183 Trp Glu His Val Asn Ala Ile Gln Glu Ala Arg Arg Leu Leu Asn Leu
184          565          570          575
185 Ser Arg Asp Thr Ala Ala Glu Met Asn Glu Thr Val Glu Val Ile Ser
186          580          585          590
187 Glu Met Phe Asp Leu Gln Glu Pro Thr Cys Leu Gln Thr Arg Leu Glu
188          595          600          605
189 Leu Tyr Lys Gln Gly Leu Arg Gly Ser Leu Thr Lys Leu Lys Gly Pro
190          610          615          620
191 Leu Thr Met Met Ala Ser His Tyr Lys Gln His Cys Pro Pro Thr Pro
192 625          630          635          640
193 Glu Thr Ser Cys Ala Thr Gln Ile Ile Thr Phe Glu Ser Phe Lys Glu
194          645          650          655
195 Asn Leu Lys Asp Phe Leu Leu Val Ile Pro Phe Asp Cys Trp Glu Pro
196          660          665          670
197 Val Gln Glu Gly Ala Pro Pro Pro Pro Ala Ala Ala His His His His
198          675          680          685
199 His His
200          690
202 <210> SEQ ID NO: 3
203 <211> LENGTH: 564
204 <212> TYPE: PRT
205 <213> ORGANISM: Artificial Sequence
207 <220> FEATURE:
208 <223> OTHER INFORMATION: HER500* construct
210 <400> SEQUENCE: 3
211 Met Arg Ala Ala Pro Leu Leu Leu Ala Arg Ala Ala Ser Leu Ser Leu
212 1 5 10 15
213 Gly Phe Leu Phe Leu Leu Phe Phe Trp Leu Asp Arg Ser Val Leu Ala

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214		20		25		30
215	Lys	Glu	Leu	Ala	Arg	Gly
216		35		40		45
217	Asp	Met	Lys	Leu	Arg	Leu
218		50		55		60
219	Leu	Arg	His	Leu	Tyr	Gln
220	65			70		75
221	Leu	Thr	Tyr	Leu	Pro	Thr
222				85		90
223	Gln	Glu	Val	Gln	Gly	Tyr
224				100		105
225	Val	Pro	Leu	Gln	Arg	Leu
226				115		120
227	Asp	Asn	Tyr	Ala	Leu	Ala
228				130		135
229	Thr	Thr	Pro	Val	Thr	Gly
230	145					150
231	Leu	Arg	Ser	Leu	Thr	Glu
232						165
233	Asn	Pro	Gln	Leu	Cys	Tyr
234						180
235	His	Lys	Asn	Asn	Gln	Leu
236						195
237	Arg	Ala	Cys	His	Pro	Cys
238						210
239	Gly	Glu	Ser	Ser	Glu	Asp
240	225					230
241	Gly	Gly	Cys	Ala	Arg	Cys
242						245
243	Glu	Gln	Cys	Ala	Ala	Gly
244						260
245	Ala	Cys	Leu	His	Phe	Asn
246						275
247	Ala	Leu	Val	Thr	Tyr	Asn
248						290
249	Glu	Gly	Arg	Tyr	Thr	Phe
250	305					310
251	Asn	Tyr	Leu	Ser	Thr	Asp
252						325
253	Lys	Leu	Gly	Ala	Gly	Gly
254						340
255	Thr	Arg	Ser	Gly	Gly	Gly
256						355
257	Glu	Glu	Ala	Pro	Arg	Ser
258						370
259	Asp	Val	Phe	Asp	Gly	Asp
260	385					390
261	Ser	Leu	Pro	Thr	His	Asp
262						405

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/821,883

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TIME: 13:12:42

Input Set : A:\7636-0022.30-SEQLIST.txt

Output Set: N:\CRF3\04232001\I821883.raw

L:14 M:270 C: Current Application Number differs, Replaced Current Application Number

L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date